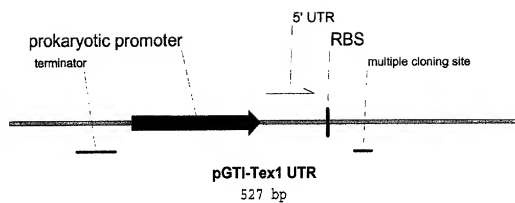
**Fig. 1**

**Fig. 2**

GGCCGGGGCCCCGCCCTTTGGGCGGGGCCTCCCCCAAGGAGGGCCG

Fig. 3

GCATGCTTATCTCGAGACTGGCAGTTCAATAGAGATATTGTATGCCTGCAG

Fig. 4

AAAGGGA

Fig. 5

GAAGGAGG

Fig. 6

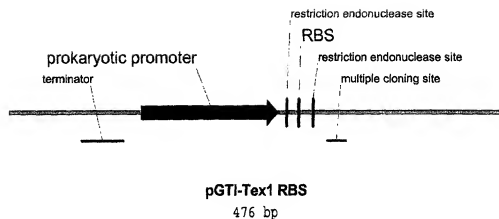


Fig. 7

GGAGGAGGCC	AAGAAGCTTT	TGGAGGGGAA	GCCCGTCTAC	ATGTACCCCA	CGTCCATTGA	60
GGCGGCCAAG	GCCATCGTGG	CCATGGTGGG	AGGTGCGGCG	TGATCCTGGT	GAACCGGAG	120
ACCCGCGTCC	TGGTCCAGGG	CATCACCGGC	CGGGAGGGGC	AGTTCCACAC	CAAGCAGATG	180
CTGGACTACG	GCACCAAGAT	CGTCGCCGGG	GTCAACCCCG	GCAAAGGGGG	AACGGAGGTC	240
CTAGGGGTCC	CCGTCTACGA	CACGGTGAAG	GAGGCGGTGG	CGCACCACGA	GGTGGACGCC	300
TCCATCATCT	TCGTGCCCGC	CCCGGCCGCG	GCGGACGCCG	CCCTGGAAGC	GGCCACGCC	360
GGGATCCCC	TCACTGTCCT	CATCACCGAG	GGCATCCCCA	CCCTGGACAT	GGTGCGGGCG	420
GTGGAGGAGA	TCAAGGCCCT	GGGAAGCCGC	CTCATCGGGG	GGAAGTCCCC	GGGGATCATC	480
AGCGCCGAGG	AGACCAAGAT	CGGGATCATG	CCCGGCCACG	TCITCAAGCG	GGGCCGGGTG	540
GGGATCATCA	GCCGCTCCGG	CACCTCACC	TACGAGGCCG	CAGCCGCCCT	TTCACAGGCG	600
GGGCTCGGCA	CCACCACCAC	GGTGGGGATC	GGGGGCGACC	CCGTCATCGG	CACCACCTTC	660
AAGGACCTCC	TCCCCCTCTT	CAACGAGGAC	CCGGAGACGG	AGGCCGTGGT	CCTCATCGGG	720
GAGATCGGCG	GCTCCGACGA	GGAGGAGGCG	GCGGCTTGGG	TGAAGGACCA	CA	772

Fig. 8